

0961201-092403  
MDEADRRLLRRCLRLVEELQVDQLWDVLLSRELFRPHMIEDIQRAGSGSRRDQA  
RQLIIDLETRGSQAL  
PLFISCLEDTGQDMLASFLRTNRQAGKLSKPTLENLTPVVLRLPEIRKPEVLRPETPR  
PVDIGSGGGFGDVG  
ALESRLRGNADLAYILSMEPCGHCLIINNPNFCRESGLRTRTGSNIDCEKLRRRFSSL  
HFMVEVKGDLTAK  
KMVLALLELARQDHGALDCCVVILSHGCQASHLQFPGAVYGTGCPVSVEKIVNI  
FNGTSCPSLGGKPK  
LFFIQACGGEQKDHGFEVASTSPEDESPGSNPEPDATPFQEGLRTFDQLDAISSLP  
TPSDIFVSYSTFPG  
FVSWRDPKSGSWYVETLDDIFEQWAHSEDLQSLLLRVANAVSVKGIYKQMPGCFN  
FLRKKLFFKTS

**FIG. 1**

1 GCCATGGACG AAGCGGATCG GCGGCTCCTG CGGCGGTGCC GGCTGCGGCT

51 GGTGGAAGAG CTGCAGGTGG ACCAGCTCTG GGACGTCCTG CTGAGCCGCG

101 AGCTGTTTCAG GCCCCATATG ATCGAGGACA TCCAGCGGGC AGGCTCTGGA

151 TCTCGGCGGG ATCAGGCCAG GCAGCTGATC ATAGATCTGG AGACTCGAGG

201 GAGTCAGGCT CTTCCITTTGT TCATCTCCTG CTTAGAGGAC ACAGGCCAGG

251 ACATGCTGGC TTCGTTTCTG CGAACTAACA GGCAAGCAGG AAAGTTGTCTG

301 AAGCCAACCC TAGAAAACCT TACCCAGTG GTGCTCAGAC CAGAGATTCTG

351 CAAACCAGAG GTTCTCAGAC CGGAAACACC CAGACCAGTG GACATTGGTT

401 CTGGAGGATT CGGTGATGTC GGTGCTCTTG AGAGTTTGAG GGGAAATGCA

451 GATTTGGCTT ACATCCTGAG CATGGAGCCC TGTGGCCACT GCCTCATTAT

501 CAACAATGTG AACTTCTGCC GTGAGTCCGG GCTCCGCACC CGCACTGGCT

551 CCAACATCGA CTGTGAGAAG TTGCGGCGTC GCTTCTCCTC GCTGCATTTC

601 ATGGTGGAGG TGAAGGGCGA CCTGACTGCC AAGAAAATGG TGCTGGCTTT

651 GCTGGAGCTG GCGCGGCAGG ACCACGGTGC TCTGGACTGC TCGTGGTGG

701 TCATTCTCTC TCACGGCTGT CAGGCCAGCC ACCTGCAGTT CCCAGGGGCT

FIG. 2A

751 GTCTACGGCA CAGATGGATG CCCTGTGTCG GTCGAGAAGA TTGTGAACAT  
801 CTTCAATGGG ACCAGCTGCC CCAGCCTGGG AGGGAAGCCC AAGCTCTTTT  
851 TCATCCAGGC CTGTGGTGGG GAGCAGAAAG ACCATGGGTT TGAGGTGGCC  
901 TCCACTTCCC CTGAAGACGA GTCCCCTGGC AGTAACCCCG AGCCAGATGC  
951 CACCCCGTTC CAGGAAGGTT TGAGGACCTT CGACCAGCTG GACGCCATAT  
1001 CTAGTTTGCC CACACCCAGT GACATCTTTG TGTCTACTC TACTTTCCCA  
1051 GGTTTTGTTT CCTGGAGGGA CCCCAAGAGT GGCTCCTGGT ACGTTGAGAC  
1101 CCTGGACGAC ATCTTTGAGC AGTGGGCTCA CTCTGAAGAC CTGCAGTCCC  
1151 TCCTGCTTAG GGTCGCTAAT GCTGTTTCGG TGAAAGGGAT TTATAAACAG  
1201 ATGCCTGGTT GCTTTAATTT CCTCCGAAA AAACTTTTCT TTAAAACATC  
1251 ATAAGGCCAG GGCCCCTCAC CCTGCCTTAT CTTGCACCCC AAAGCTTTCC  
1301 TGCCCCAGGC CTGAAAGAGG CTGAGGCCTG GACTTTCCTG CAACTCAAGG  
1351 ACTTTGNAGC CGGCACAGGG TCTGCTCTTT CTCTGCCAGT GACAGACAGG  
1401 CTCTTAGCAG CTTCCAGATT GACGACAAGT GCTGAACAGT GGAGGAAGAG  
1451 GGACAGATGA ATGCCGTGGA TTGCACGTGG NCTCTTGAGC AGTGGCTGGT

FIG. 2B

1501 CCAGGGCTAG TGA CT TGGTG TCCCATGATC CCTGTGTTGG TCTCTAGGAG

1551 CAGGGATTAA CCTCTGCACT ACTGACAT

## FIG. 2C

CTGACTGCCAAGAAAATGGTGCTGGCTTTGCTGGAGCTGG 40  
CGCGGCAGGACCACGGTGCTCTGGACTGCTGCGTGGTGGT 80  
CATTCTCTCTCACGGCTGTCAGGCCAGCCACCTGCAGTTC 120  
CCAGGGGCTGTCTACGGCACAGATGGATGCCCTGTGTCCG 160  
TCGAAAAGATTGTGAACATCTTCAATGGGACCAGCTGCCC 200  
CAGCCTGGGAGGGAAGCCCAAGCTCTTTTTTCATCCAGGCC 240  
TGTGGTGGGGAGCAGAAAGACCATGGGTTTGAGGTGGCCT 280  
CCACTTCCCCTGAAGACGAGTCCCCTGGCAGTAACCCCGA 320  
GCCAGATGCCACCCCGTTCCAGGAAGGTTTGAGGACCTTC 360  
GACCAGCTGGACGCCATATCTAGTTTGCCACACCCAGTG 400  
ACATCTTTGTGTCCTACTCTACTTTCCCAGGTTTTGTTTC 440  
CTGGAGGGACCCCAAGAGTGGCTCCTGGTACGTTGAGACC 480  
CTGGACGACATCTTTGAGCAGTGGGCTCACTCTGAAGACC 520  
TGCAGTCCCTCCTGCTTAGGGTCGCTAATGCTGTTTCGGT 560  
GAAAGGGATTTATAAACAGATGCCTGGTTGCTTTAATTTTC 600  
CTCCGGAAAAAACTTTTCTTTTAAAACATCATAAGGCAG 639

## FIG. 3

0961201-092401  
204250-2021950

MVLALLELARQDHGALDCCV 20  
VVILSHGCQASHLQFPGAVY 40  
GTDGCPVSVEKIVNIFNGTS 60  
CPSLGGKPKLFFIQACGGEQ 80  
KDHGFEVASTSPEDESPGSN 100  
PEPDATPFQEGLRTFDQLDA 120  
ISSLPTPSDIFVSYSTFPGF 140  
VSWRDPKSGSWYVETLDDIF 160  
EQWAHSEDLQSLLLRVANAV 180  
SVKGIYKQMPGCFNFLRKKL 200  
FFM 203

**FIG. 4**

ICE/CED3  
GENE FAMILY:

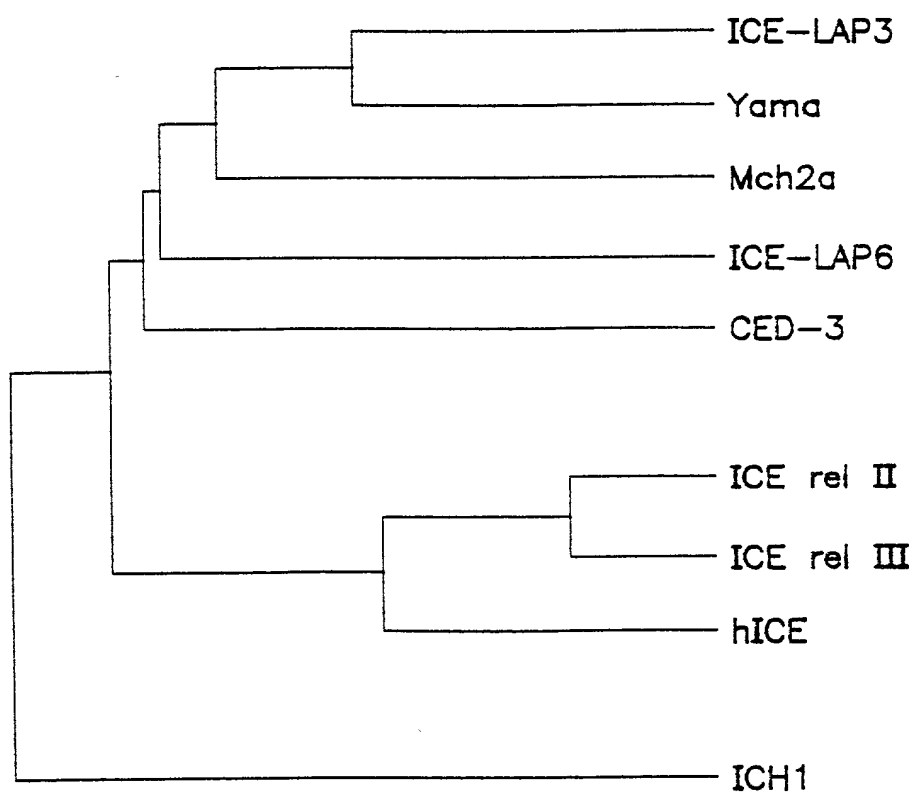


FIG. 5

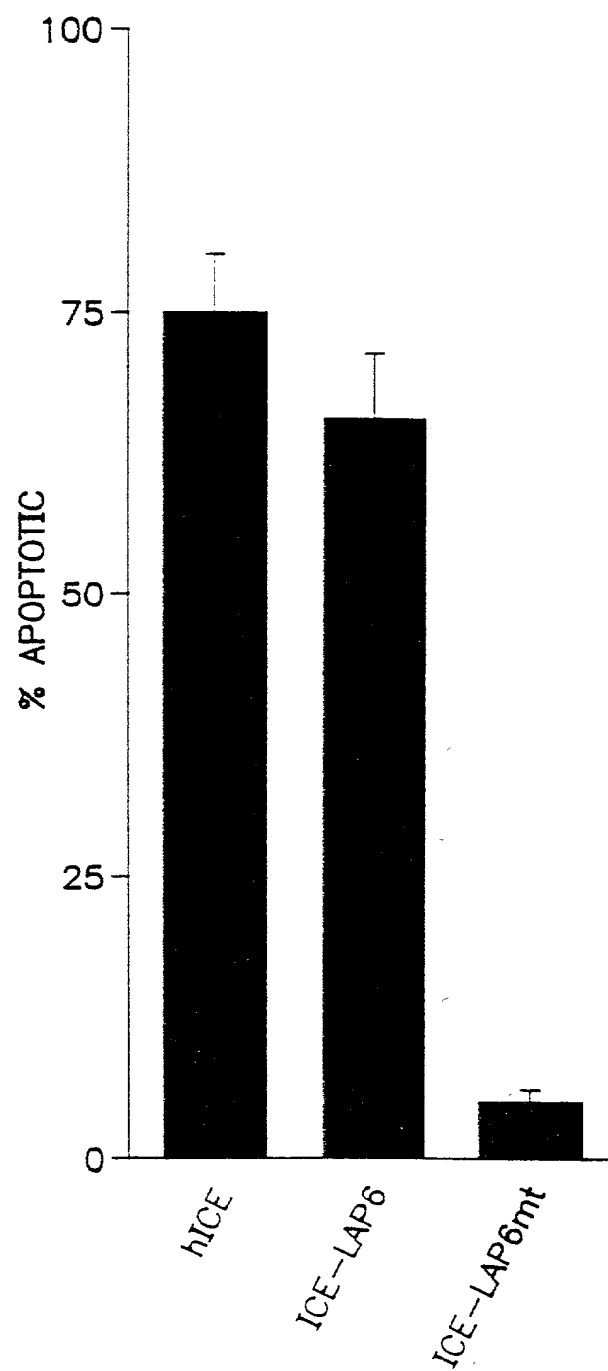


FIG. 6